

Author Correction: TDP-43 loss and ALS-risk SNPs drive mis-splicing and depletion of UNC13A


<https://doi.org/10.1038/s41586-024-07577-9>

Published online: 18 June 2024

Correction to: *Nature* <https://doi.org/10.1038/s41586-022-04436-3>

Published online 23 February 2022

Open access

 Check for updates

Anna-Leigh Brown, Oscar G. Wilkins, Matthew J. Keuss, Sarah E. Hill, Matteo Zanovello, Weaverly Colleen Lee, Alexander Bampton, Flora C. Y. Lee, Laura Masino, Yue A. Qi, Sam Bryce-Smith, Ariana Gatt, Martina Hallegger, Delphine Fagegaltier, Hemali Phatnani, NYGC ALS Consortium, Jia Newcombe, Emil K. Gustavsson, Sahba Seddighi, Joel F. Reyes, Steven L. Coon, Daniel Ramos, Giampietro Schiavo, Elizabeth M. C. Fisher, Towfique Raj, Maria Secrier, Tammaryn Lashley, Jernej Ule, Emanuele Buratti, Jack Humphrey, Michael E. Ward & Pietro Fratta

The version of the article initially published referred to one, cell line as SK-N-DZ. This line has since been conclusively identified as SK-N-BE(2). The HTML and PDF versions of the article have been updated to reflect this.



Open Access This article is licensed under a Creative Commons Attribution 4.0 International License, which permits use, sharing, adaptation, distribution and reproduction in any medium or format, as long as you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons licence, and indicate if changes were made. The images or other third party material in this article are included in the article's Creative Commons licence, unless indicated otherwise in a credit line to the material. If material is not included in the article's Creative Commons licence and your intended use is not permitted by statutory regulation or exceeds the permitted use, you will need to obtain permission directly from the copyright holder. To view a copy of this licence, visit <http://creativecommons.org/licenses/by/4.0/>.

© The Author(s) 2024